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The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 101643, 752
Source: 0126
Date Processed by STIC: 8-25-03

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS. PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- Hand Carry directly to:
 U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name,
 Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
 - U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
- Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1803-Mallroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 04/24/2003

ERROR DETECTED	SUCCESTED CORRECTION SERIAL NUMBER: 10 1043, 702
ATTN: NEW RULES CASES:	Please disregard english "Alpha" headers, which were inserted by Pto Software
1Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.
3Misaligned Amino Numbering	The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
4Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
5Variable Length	Sequence(s)contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6PatentIn 2.0 "bug"	A "bug" in Patentin version 2.0 has caused the <220><223> section to be missing from amino acid sequences(s)
7Skipped Sequences (OLD RULES)	Sequence(s)missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (1) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
	(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped
	Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to Include the skipped sequences.
8 Skipped Sequences (NEW RULES)	Sequence(s) missing. If Intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000
9Uze of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220> <223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa; and which residue n or Xaa represents.
10Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220><223> section is required when <213> response is Unknown or is Artificial Sequence
11 X Use of <220>	Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
12Patentin 2.0 ~ "bug"	Please do not use "Copy to Disk" function of Patentin version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Imstead, please use "File Manager" or any other manual means to copy file to floppy disk.
13Misuse of n	n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.
	AMC/ASH - Rintechnology Systems Branch - 02/21/2001



DATE: 08/25/2003

TIME: 14:10:45

OIPE

Input Set : A:\LS5-001.ST25.txt Output Set: N:\CRF4\08252003\J643752.raw 3 <110> APPLICANT: Liu, David R. Gartner , Zev J. Doyon, Jeffrey B. 5 Calderone , Christopher T. 7 Kanan, Matthew W. Li, Xiaoyu 9 Snyder, Thomas M. Rosenbaum, Daniel M. 10 12 <120> TITLE OF INVENTION: Evolving New Molecular Function 14 <130> FILE REFERENCE: LS5-001 C--> 16 <140> CURRENT APPLICATION NUMBER: US/10/643,752 Does Not Comply C--> 16 <141> CURRENT FILING DATE: 2003-08-19 Corrected Diskette Needed 16 <150> PRIOR APPLICATION NUMBER: US 60/404,395 17 <151> PRIOR FILING DATE: 2002-08-19 19 <150> PRIOR APPLICATION NUMBER: US 60/419,667 20 <151> PRIOR FILING DATE: 2002-10-18 22 <150> PRIOR APPLICATION NUMBER: US 60/432,812 23 <151> PRIOR FILING DATE: 2002-12-11 25 <150> PRIOR APPLICATION NUMBER: US 60/444,770 26 <151> PRIOR FILING DATE: 2003-02-04 28 <150> PRIOR APPLICATION NUMBER: US 60/457,789 29 <151> PRIOR FILING DATE: 2003-03-26 31 <150> PRIOR APPLICATION NUMBER: US 60/469,866 32 <151> PRIOR FILING DATE: 2003-05-12 34 <150> PRIOR APPLICATION NUMBER: US 60/479,494 35 <151> PRIOR FILING DATE: 2003-06-18 37 <160> NUMBER OF SEQ ID NOS: 125 39 <170> SOFTWARE: PatentIn version 3.1 41 <210> SEQ ID NO: 1 42 <211> LENGTH: 64 43 <212> TYPE: DNA 44 <213> ORGANISM: Artificial Sequence 46 <220> FEATURE: 47 <223> OTHER INFORMATION: Template Encoding Parent Molecule 1 49 <400> SEQUENCE: 1 60 50 cgagcagcac cagcgcactc cgcctggatc cgcccgggt gcacgcgact cctacgggct 64 52 ccaa 55 <210> SEQ ID NO: 2 56 <211> LENGTH: 64 57 <212> TYPE: DNA 58 <213> ORGANISM: Artificial Sequence 60 <220> FEATURE: 61 <223> OTHER INFORMATION: Template Encoding Parent Molecule 2

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/643,752

RAW SEQUENCE LISTING DATE: 08/25/2003
PATENT APPLICATION: US/10/643,752 TIME: 14:10:45

Input Set : A:\LS5-001.ST25.txt

Output Set: N:\CRF4\08252003\J643752.raw

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 66 ccaa
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 71 <212> TYPE: DNA
 72 <213> ORGANISM: Artificial Sequence
 74 <220> FEATURE:
 75 <223> OTHER INFORMATION: Recombined Daughter Template
 77 <400> SEQUENCE: 3
 78 cgagcagcac cagcgcactc cgcctgggga tgccccgggt gggcgcgact cctacgggct
 80 ccaa
 83 <210> SEQ ID NO: 4
 84 <211> LENGTH: 64
 85 <212> TYPE: DNA
 86 <213> ORGANISM: Artificial Sequence
 88 <220> FEATURE:
 89 <223> OTHER INFORMATION: Recombined Daughter Template
 91 <400> SEQUENCE: 4
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 94 ccaa
97 <210> SEQ ID NO: 5
98 <211> LENGTH: 10
99 <212> TYPE: DNA
                                              inatticient response
100 <213> ORGANISM: Artificial Sequence
                                           please sive source of sometic material.
102 <220> FEATURE:
103 <223> OTHER INFORMATION: Reagent
105 <400> SEQUENCE: 5
                                             (sæitem 11 on error 10
summary report.
106 aattcgtacc
109 <210> SEQ ID NO: 6
. 110 <211> LENGTH: 11
111 <212> TYPE: DNA
112 <213> ORGANISM: Artificial Sequence
114 <220> FEATURE:
115 <223> OTHER INFORMATION: Template E
117 <400 SEQUENCE: 6
118 tggtacgaat t
                                                                            11
121 <210> SEQ ID NO: 7
122 <211> LENGTH: 31
123 <212> TYPE: DNA
124 <213> ORGANISM: Artificial Sequence
126 <220> FEATURE:
127 <223> OTHER INFORMATION: Template H
129 <400> SEQUENCE: 7
130 tcgcgagcgt acgctcgcga tggtacgaat t
                                                                            31
133 <210> SEQ ID NO: 8
134 <211> LENGTH: 20
135 <212> TYPE: DNA
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RAW SEQUENCE LISTING DATE: 08/25/2003
PATENT APPLICATION: US/10/643,752 TIME: 14:10:45

Input Set : A:\LS5-001.ST25.txt

Output Set: N:\CRF4\08252003\J643752.raw

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    146 <211> LENGTH: 10
    147 <212> TYPE: DNA
    148 <213> ORGANISM: Artificial Sequence
    150 <220> FEATURE:
    151 <223> OTHER INFORMATION Reagent
    153 <400> SEQUENCE: 9
                                                                                10
    154 cccgagtcga
    157 <210> SEQ ID NO: 10
    158 <211> LENGTH: 50
    159 <212> TYPE: DNA
    160 <213> ORGANISM: Artificial Sequence
    162 <220> FEATURE:
    163 <223> OTHER INFORMATION: Template
    165 <400> SEQUENCE: 10
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    169 <210> SEQ ID NO: 11
    170 <211> LENGTH: 50
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    172 <213> ORGANISM: Artificial Sequence
    174 <220> FEATURE:
    175 <223> OTHER INFORMATION: Template
    177 <220> FEATURE:
    178 <221> NAME/KEY: misc feature
    179 <222> LOCATION: (17)..(17)
    180 <223> OTHER INFORMATION: N is A, C, T or G
    183 <220> FEATURE:
    184 <221> NAME/KEY: misc_feature
    185 <222> LOCATION: (19)..(19)
    186 <223> OTHER INFORMATION: N is A, C, T or G
    189 <220> FEATURE:
    190 <221> NAME/KEY: misc_feature
    191 <222> LOCATION: (21)..(21)
    192 <223> OTHER INFORMATION: N is A, C, T or G
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    196 <221> NAME/KEY: misc_feature
    197 <222> LOCATION: (23)..(24)
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                                                                                50
W--> 202 tggtgoggag cogcognona nonngatace acotecgage cgaggageeg
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     206 <211> LENGTH: 10
    207 <212> TYPE: DNA
    208 <213> ORGANISM: Artificial Sequence
     210 <220> FEATURE:
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RAW SEQUENCE LISTING DATE: 08/25/2003 PATENT APPLICATION: US/10/643,752 TIME: 14:10:45

Input Set : A:\LS5-001.ST25.txt

Output Set: N:\CRF4\08252003\J643752.raw

```
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     213 <400> SEQUENCE: 12
     214 cacccgtcac
                                                                                10
     217 <210> SEQ ID NO: 13
     218 <211> LENGTH: 10
     219 <212> TYPE: DNA
     220 <213> ORGANISM: Artificial Sequence
     222 <220> FEATURE:
     223 <223> OTHER INFORMATION Reagent
     225 <220> FEATURE:
     226 <221> NAME/KEY: misc_feature
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    228 <223> OTHER INFORMATION: N is A, T, C or G
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    233 <222> LOCATION: (5)..(5)
    234 <223> OTHER INFORMATION: N is A, T, C or G
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    246 <223> OTHER INFORMATION: N is A, T, C or G
    249 <400> SEQUENCE: 13
W--> 250 cnngntngnc
                                                                                10
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    256 <213> ORGANISM: Artificial Sequence
    258 <220> FEATURE:
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    262 tggtacgaat t
                                                                                11
    265 <210> SEQ ID NO: 15
    266 <211> LENGTH: 17
    267 <212> TYPE: DNA
    268 <213> ORGANISM: Artificial Sequence
    270 <220> FEATURE:
    271 <223> OTHER INFORMATION: Template 2a-2c
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                                                                                17
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    278 <211> LENGTH: 23
    279 <212> TYPE: DNA
   280 <213> ORGANISM: Artificial Sequence
    282 <220> FEATURE:
    283 <223> OTHER INFORMATION: Template 3a-3c
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RAW SEQUENCE LISTING DATE: 08/25/2003 PATENT APPLICATION: US/10/643,752 TIME: 14:10:45

Input Set : A:\LS5-001.ST25.txt

Output Set: N:\CRF4\08252003\J643752.raw

	285	<400> SEQUENCE: 16	
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	289	<210> SEQ ID NO: 17	
	290	<211> LENGTH: 14	
		<212> TYPE: DNA	
		<213> ORGANISM: Artificial Sequence	
		<220> FEATURE:	
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		<400> SEQUENCE: 17	
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	301	<210> SEQ ID NO: 18	
		<211> LENGTH: 16	
	303	<212> TYPE: DNA	
	304	<213> ORGANISM: Artificial Sequence	
	306	<220> FEATURE:	
	307	<223> OTHER INFORMATION: Reagent 5a-5c	
	309	<400> SEQUENCE: 18	
	310	ctcagctctc tcgtta	16
	313	<210> SEQ ID NO: 19	
	314	<211> LENGTH: 18	
	315	<212> TYPE: DNA	
	316	<213> ORGANISM: Artificial Sequence	
	318	<220> FEATURE:	
	319	<223> OTHER INFORMATION: Reagent 6a-6c	
		<400> SEQUENCE: 19	
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		<210> SEQ ID NO: 20	
		<211> LENGTH: 11	
		<212> TYPE: DNA	
•		<213> ORGANISM: Artificial Sequence	
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	-	<223> OTHER INFORMATION: Template 15	
		<400> SEQUENCE: 20	
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		<211> LENGTH: 11	
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		<400> SEQUENCE: 21	
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		<210> SEQ ID NO: 22	
		<211> LENGTH: 11	
		<211> DENGTH. IT <212> TYPE: DNA	
		<213> ORGANISM: Artificial Sequence	
		<220> FEATURE:	
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		<400> SEQUENCE: 22	
	33 I	ZADON GENOEILE: 22	

The type of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 08/25/2003

PATENT APPLICATION: US/10/643,752 TIME: 14:10:46

Input Set : A:\L85-001.ST25.txt

Output Set: N:\CRF4\08252003\J643752.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:11; N Pos. 17,19,21,23,24

Seq#:13; N Pos: 2,3,5,7,9

Seq#:31; N Pos. 11,12,13,14,15,16,22,23,24,25,26,27,33,34,35,36,37,38

Seq#:34; N Pos. 17,18,19,20,21,22,23,24,25,26,27,28,29,30,31,32,33,34,35,36

Seq#:60; N Pos. 14

VERIFICATION SUMMARY

DATE: 08/25/2003 PATENT APPLICATION: US/10/643,752 TIME: 14:10:46

Input Set : A:\LS5-001.ST25.txt

Output Set: N:\CRF4\08252003\J643752.raw

L:16 M:270 C: Current Application Number differs, Replaced Current Application No L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:202 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0 L:250 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0 L:484 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:0 L:526 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:0 L:844 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:60 after pos.:0